13.		in Cos and		*	1/15			<u> </u>			j) s
ergleichsekrivensusprolenasi	TRADE (umop 1 - dn 81) dn	up (17-2)	up (9-1)		up (17-0)	up (17-0)	up (22-2)	up (17-1)	up (25-0)	up (11-0)	up (10-2)
Vergleich Endometriose IV. Vergleich Endometriose IV. Vergleich Endometriose IV. Vergleich IV. Versust Vormali(prolifichtese)	down (4 up -12 down)	nc (13-13)	nc (6-2)		nc (8-10)	down (0-8)	down (4-13)	nc (9-10)	nc (0-0)	nc (3-3)	nc (9-10)
Vergleich Endometriose 45 (III)	down (0 up - 16 down)	down (1-15)	down (0-14)		down (0-13)	down (0-13)	down (1-13)	down (1-13)	down (0-12)	down (0-12)	down (1-12)
The state of the s		S37730, insulin-like growth factor binding protein-2	U40271, Human transmembrane receptor precursor	(PTK7)	M21574, platelet-derived growth factor receptor alpha	1 22548 collagen type XVIII alpha 1 (COL18A1)	1400482 cubtilisin-like protein (PACE4)	726653. Jaminin M chain (merosin)	Magged 1177846, Elastin	x05610, type IV collagen alpha -2 chain	X67325, p27 interferon alpha-inducible gene

Abbildung l

[Key to Table:]

Datenbank-Nr., Name = Data Bank No., Name

Vergleich Endometriose versus Normal (sekr. Phase) = Comparison of Endometriosis versus Normal (Secr. Phase)

Vergleich Endometriose versus Normal (prol. Phase) = Comparison of Endometriosis versus Normal (Prol. Phase)

Vergleich sekr. versus prol. Phase (Endometrium) = Comparison of Secr. versus Prol. Phase (Endometrium)

[Key to Table:]

Datenbank-Nr., Name = Data Bank No., Name
Vergleich Endometriose versus Normal (sekr. Phase) = Comparison
 of Endometriosis versus Normal (Secr. Phase)
Vergleich Endometriose versus Normal (prol. Phase) = Comparison
 of Endometriosis versus Normal (Prol. Phase)
Vergleich sekr. versus prol. Phase (Endometrium) = Comparison
 of Secr. versus Prol. Phase (Endometrium)

down (0-11) down (1-11) down (0-10)	Datenbank-Nr., Name	Vergleich Endometriose versus Normal (sekr. Phase)	Vergleich Endometriose versus Normal (prol. Phase)	Vergleich selcr. versus prol. Phase (Endometrium)
de dehydrogenase 6 down (1-11) en down (0-10)	D42073, reticulocalbin	down (0-11)	nc (8-5)	up (11-2)
en down (1-11) down (0-10)	U07919, aldehyde dehydrogenase 6	down (1-11)	ne (13-9)	up (22-0)
down (0-10)	U81607, gravin	down (1-11)	nc (8-7)	up (18-1)
() [] amop	M30269, nidogen	down (0-10)	nc (8-14)	up (15-3)
(o1-1) II won	D42408, phospholipase C Epsilon	down (1-10)	nc (12-14)	up (25-0)

Abbildung 1

Abbildung 2		
Seq.IDNO	Name	Proteinsequenz
	Fibronektin	MLAGAGAGIL LIANQCLGTA VPSTGASKSK RQAQQMVQPQ SPVAVSQSKP GCYDNGKHYQ INQOWBRTYL GNALVCTCYG GSRGFNCESK PEAEETCFDK YTGNTYRVGD TYERPKDSMI WDCTCLGAGR GRISCTIANR CHEGOGSYKI GDTWRPHET GGYMLECVCL GNGKGBWTCK PIAEKCFDHA AGTSYVVGET WEKPYGGWMM CHEGOGSYKI GDTWRPHET GGYMLECVCL GNGKGBWTCK PIAEKCFDHA AGTSYVVGET WEKPYGGWMM SGSGPFTDVR AAVYQPQPHP GPPPYGHCYT BGGVYSVGW GHLKTQGIK GHLKTQGIKS GGSGPFTDVR AAVYQPQPHP GPPPYGHCYT BGGVYSVGW GHLKTQGIKG MLCTCLGNGV SCGFTAVTQT SGSGPFTDVR AAVYQPQPHP GPPPYGHCYT BGGVYSVGW GHLKTQGIKG MLCTCLGNGV SCGFTAVTQT GGMSNGEPC VLPFTXNGRT FYSCTTEGRQ DGHLNCSTTS NYEDDOKXSF CTDHTVLVQT GGGNSNGALC YGGNSNGEPC VLPFTXNGRT FYSCTTEGRQ DGHLNSTTKG LKPGVYVEGQ LISIQQYGHQ EVTREPPTTT NAPQPSHISK YILKWRPKNS VGRWKEATIP GHLNSYTIKG LKPGVVYEGQ LISIQQYGHQ EVTREPPTTT NAPQPSHISK YILKWRPKNS VGRWKEATIP GHLNSYTIKG LKPGVVYEGQ LISIQQYGHQ EVTREPPTTT NAPQPSHISK YILKWRPKNS VGRWKEATIP GHLNSYTIKG LKPGVVYEGQ LISIQQYGHQ EVTREPPTTT NAPQPSHISK YILKWRPKNS VGRWKEATIP GHLNSYTIKG LKPGVYYEGG LISIQQYGHQ EVTREPPTTT NAPQPSHISK YILKWRPKNS VGRSEDGEGG LILSTSGTTA PARSTYNFG LFGSGVYSTOV PROLEFSRAT FARSVILSDL PROSTYKEVY YRLVYSPSVE GSSTELMLPR TANGYTLSDL QPGVQYNITI YAVENOGEST PVVIQQFTG TPRSDPTYD TATSVNIPDL LPGRKYTUMY YGISEDGEGG LILSTSGTTA PARSTYNFWY TTTVITMTPA PRIGFKLGVR YPLRNLQPAS EYTVSLIVARTY GNGESPKATG VFTLLQPGSS IPPVNTEVTF TTTVITMTPA PRIGFKLGVR YPLRNLQPAS EYTVSLIVAR GNGESPKATG VFTLLQPGSS IPPVNTEVTF TTTVITMTPA PRIGFKLGVR YPLRNLQPAS EYTVSLIVAR GNGESPKATG VFTLLQPGS IPPVNTEVTF TTTVITMTPA PRIGFKLGVR GTERVVSVSS VYEGHESTPL GRACKTGLDS GTERVVSVSS VYEGHESTPL GRACKTGLDS GTERVVSVSS VYEGHESTPL GRACKTGLDS GTERVVSVSS VYEGHESTPL GRACKTGLDS TATSTYNTHIAP PRAGGAGGNS LEEVVHANDS PROLEVINIA PRATITGYRI THPPREFSGR GTERVVSVSS VYEGHESTPL GRACKTGLDS TATSTYNTHAP PRAGGAGGNS LEEVVHANDS PROLEVINIA PRATITGYRI THPPREFSGR GTERVVPSVS VYEGHESTPL GRACKTGLDS TATSTYNTHAP PRAGGAGGNS LAGGESPLL INGGCSTVSDV PROLESTSTSDNDAVITNILLISMDA
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[Key to Table:]

Proteinsequenz = Protein Sequence Fibronektin = fibronectin ; ::

Abbildung 2

Seq.IDNO	Name	Proteinsequenz
		EIDKPSQMQV TVGETGGNSP VQEFTVPGSK STATISGLKP GVDYTITVYA VTGRGDSPAS SKPISINYRT EIDKPSQMQV TDVQDNSISV KWLPSSSPVT GYRVTTPKN GPGPTKTKTA GPDQTEMTIE GLQPTVEYVV SVYAQNPSGE SQPLVQTAVT NIDRPKGLAF TDVDVDSIKI AMESPQGQVS RYRVTYSSPE DGIHELFPAP GYRVAVTPKE KTGFWKEINL APDSSSVVVS GLMVATKYEV SVYALKDTLT SRPAQGVVTT LENVSPPRRA GYRVRVTPKE KTGFWKEINL APDSSSVVVS GLMVATKYEV SVYALKDTLT SRPAQGVVTT LENVSPPRRA RSSPVVIDAS TAIDAPSNLR FLATTPNSLL VSWQPPRARI TGYIIKYEKP GSPPREVVPR PRPGVTEATI TGLEPGTEYT IYVIALKANQ KSEPLIGRKK TDELPQLVTL PHPNLHGPEI LDVPSTVQKT PFVTHPGYDT GNGIQLPGTS GQQPSVGQQM IFEEHGFRRT TPPTTATPIR HRPRPYPPNV GELQIGHIP REDVDYHLYP GNGIQLPGTS GQQPSVGQQM IFEEHGFRRT TPPTTATPIR HRPRPYPPNV GETGIGHIP REDVDYHLYP GHFRCDSSRW CHDNGVNYKI GEKWDRQGEN GQMNSCTCLG NGKGEFKCDP HEATCYDDGK TYHVGEQMQK GHFRCDSSRW CHDNGVNYKI GEKWDRQGEP SPEGTTGQSY NQYSQRYHQR TNTNVNCPIE CFMPLDVQAD REDSRB
7	Insulin-like growth factor binding protein-2	MLPRVGCPAL PLPPPPLLPL LPLLLLLGA SGGGGGRAE VLFRCPPCTP ERLAACGPPP VAPPAAVAAV AGGARMPCAE LVREPGCGCC SVCARLEGEA CGVYTPRCGQ GLRCYPHPGS ELPLQALVMG EGTCEKRRDA EYGASPEQVA DNGDDHSEGG LVENHVDSTM NMLGGGGSAG RKPLKSGMKE LAVFREKVTE QHRQMGKGGK HHLGLEEPKK LRPPPARTPC QQELDQVLER ISTMRLPDER GPLEHLYSLH IPNCDKHGLY NLKQCKMSLN GQRGECWCVN PNTGKLIQGA PTÌRGDPECH LFYNEQQEAR GVHTQRMQ

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Seq.IDNO	Name	Proteinsequenz					
	Greatmonogram	MGBARGSPAR PRRLPLLSVL LLF	LLPLLGGTQT A	AIVFIKQPSS (ODALOGRRAL I	LRCEVEAPGP V	VHVYWLLDGA
· ·	recentor PFK7	OGSSLSFAAV			EARSANASFN I	IKWIEAGPVV 1	LKHPASEAEI
		HIDGHPRPTY		DGQSNHTVSS	KERNLTLRPA C	GPEHSGLYSC (CAHSAFSQAC
		DESFARVVLA	PQDVVVARYE E	EAMFHCQFSA (OPPSSLQWLF 6	EDETPITNRS	RPPHLRRATV
		VRPRNAGIYR	CIGQGQRGPP I	IILEATLHLA	EIEDMPLFEP F	RVFTAGSEER	VTCLPPKGLP
		EPSVWWEHAG VRLPTHGRVY QKC	QKGHELVLAN I.	IAESDAGVYT	CHAANLAGOR F	RODVNITVAT	VPSWLKKPQD
. •		SQLEEGKPGY LDCLTQATPK PT	PTVVWYRNOM L	LISEDSRFEV	FENGTLRINS V	VEVYDGTWYR (CMSSTPAGSI
		KLKFTPPPQP	QOCMGFDKEA T	TVPCSATGRE	KPTIKWERAD (GSSLPEWVTD	NAGTLHFARV
		IASNGPQGQI	RAHVQLTVAV F	FITFKVEPER	TTVYQCHTAL]	LOCEAQGDPK	PLIQWKGKDR
		MHIFQNGSLV	IHDVAPEDSG R	RYTCIAGNSC	NIKHTEAPLY	VVDKPVPEES	EGPGSPPPYK
		AAVAYIIAVL	GLMFYCKKRC K	KAKRLOKOPE	GEEPEMECLN (SGDNÖTGDD	AEIQEEVALT
		KRHSTSDKMH	FPRSSLQPIT T	TLGKSEFGEV	FLAKAQGLEE (GVAETLVLVK	SLQSKDEQQQ
		GKLNHANVVR	LLGLCREAEP H	IYMVLEYVDL	HYMVLEYVDL EDLKQFLRIS	KSKDEKLKSQ	PLSTKQKVAL
			LAARNCLVSA Ç	QRQVKVSALG	LSKDVYNSEY	YHFRQAFIVAL	RWMSPEAILE
		ASGVLMWEVF	тнсемрнссо д	ADDEVLADLQ	AGKARLPOPE	GCPSKLYRLM	ÓRCWALSPKD
		RPSFSEIASA LGDSTVDSKP				:	
					·		
	Plarelet-derived	MGTSHPAFLV LGCLLTGLSL II	ILCQLSLPSI I	LPNENEKVVQ	LNSSFSLRCF	GESEVSWQYP	MSEEESSDVE
	growth factor	IRNEENNSGL FVTVLEVSSA	SAAHŢĢLYTC	YYNHTQTEEN	ELEGRHIYIY	VPDPDVAFVP	LGMTDYLVIV
	receptor alpha	EDDDSAIIPC RTTDPETPVT LA	LHNSEGVVPA :	SYDSRQGFNG	TFTVGPYICE	ATVKGKKFQT	IPFNVYALKA
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Seq.IDNO	Name	Proteinsequenz
		1
		AARQATREVK EMKKVTISVH EKGFIEIKPT FSQLEAVNLH
		QEIRYRSKLK LIRAKEEDSG HYTIVAONED AVKSTIFELL
		VLVLLVIVII SLIVLVVIGK
		EYIYVDPMQL PYDSRWEFPR
		LMSELKIMTH LGPHLNIVNL LGACTKSGF1 1111E1CF13 ZETTENNON NOADESTREY QYVPMLERKE VSKYSDIQRS
		TLLDLLSFTY QVARGMEFLA SKNCVHRDLA
		SKGSTFLPVK WMAPESIFDN LYTTLSDVWS YGILLWEIFS LGGTPYFUMM
		FYHLSEIV ENLLFGUIAA
		KDWEGGLDEQ KLSADSGIII FEEDING
ď	Collagen type	GEVGADGIPG FPGLPGREGI AGPQGPKGDR
<u>.</u>	XVIII alpha l	KGNLGSKGEL GSFGFNGENG ELGSTSTST
		PGMNGLKGER GEFGEAGES CHICKEN
		GNOGPPGPKG PKGEVGFGF FGZ: 1: 2: 2: 2: 2: 2: 2: 2: 2: 2: 2: 2: 2: 2:
		PGPRGYPG1P GPRGESINGS 13:15:25
		PPGPPGIMGA SECONDING REHPHPTARP WRADDILASP PGLPEPQPYP
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Seq.IDNO	Name	Proteinsequenz
		TUBBLICATION DIVINITION FOR PERMEALES SECPLINEGAR IFSFDGKDVL RHPTWPQKSV WHGSDPNGRR
		TEN DEATHEON
	odit nioittido	MPPRAPPE PRPPERAFA TOTAAGAGGA GGAGGAGGPG FRPLAPRPMR WLLLLALPAA CSAPPPRPVY
9		
•	procein (race)	VRSDPQALYF NDPIMSNMMY LHCGDKNSRC RSEMNVQAAW KRGYTGKNVV
		SYASYDVNGN DYDPSPRYDA SNENKHGTRC AGEVAASANN SYCIVGIAYN
		SLGIRPNYID IYSASWGPDD DGKTVDGPGR LAKQAFEYGI KKGRQGLGSI
		ALALEANSOL
·		MCVAASDKRP
		I AKBLI DI SN
		DALLOSDSOMI ELSAPELEPP
		SANGSKENNIN GEOVETSRKCV SVCPLGYFGD TAARRCRRCH
		VIT CORGENA DESORNCLIC
		MEGARCYLLI LILSGGLGGV OAQRPQQRQ SQAHQQRGLF PAVLNLASNA LITTNATCGE KGPEMYCKLV
7	Laminin M Chain	INFORMACION POCRICIONS SNPNORHPIT
	(Merosin	
		GRPSADDPSP ELLEFTSARY IRLRFORIRT LNADLMMFAH KDPREIDPIV
		ISVGGMCICY GHARACPLDP ATNKSRCECE HNTCGDSCDQ CCPGFHQKPW RAGTFLTKTE CEACNCHGKA

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eq.IDNO	Name	Proteinsequenz
		1
		RKNLSLN1KG ALLGGGVCT.
		SCHCKTGFGG VSCDNCAMOS YWTYGKTODM
		KSGFFNLQED NWKGCDECFC SGVSWACGS ::::
		NAEARQALPH SIIMSAFAFI LONNEIN C
		VYLHPSEEHT NVLLLINGEST TINGTINGTON VYLHPSEEHT NVLLLINGEST TINGTINGTIF
		YPTDGSIAAA VEVCQCFFGI ISEECT
:		GFIGEFING! JEPCKI CHC.
		SCOPCOCNUM LDFStruston CKAGTERIOS
		ANVOGORCDA CINAGO I CESSO
	•	
		MUSTYGSLDF QCNVNTGQCH CHPKFSGAKC TECSRGHMNY PRCNLCDCFL
		VINCACATOOR
		TOTOLICIANT
		Vrunesi vai incolo
		Number of the state of the stat
		GPTLGTCVPC QCNGHSSLCD PETSICONCO HHTAGDFCER CALGYYGIVK
		SPACYARGID DYRCTACPRG YEGOYCERCA PGYTGSPGNP GGSCOECECD
		SI SCHIEGE KHWHAREGWE
		GALGRACDEC JULIUS INC.
		HLLSPORAFE KLICLARGINE KENLEGLOKE
		AVNEKAIKLN EILGIKDERI EIGHEN ZUN AKAMDIA
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[Key to Table:]

Seq.IDNO	Name	Proteinsequenz
		SOARSHAAOL NDSSAVLDGI LDEAKNISFN ATAAFKAYSN IKDYIDEAEK VAKEAKDLAH EATKLATGPR
		WANYKTAVAD NLLFYLGSAK FIDFLAIEMR KGKVSFLWDV GSGVGRVEYP DLTIDDSYWY RIVASRTGRN
		KEGDCKGCTV
		KWKSFTLSRI OKQANISIVD IDTNQEENIA TSSSGNNFGL DLKADDKIYF GGLPTLRNLS MKARPEVNLK
		GIILLGSGGT PAPPRRKRRQ TGQAYYVILL NRGRLEVHLS TGARTMRKIV IRPEPNLFHD GREHSVHVER
		TRGIFTVQVD ENRRYMQNLT VEQPIEVKKL FVGGAPPEFQ PSPLRNIPPF EGCIMNLVIN SVPMDFARPV
		TKVKNRLTIE

	رة د	MAGE, TARAPR PGVLLLILESI LHPSRPGGVP GAIPGGVPGG VFYPGAGLGA LGGGALGPGG KPLKPVPGGL
3		
		VPGVGLPGVY PGGVLPGARF PGVGVLPGVP TGAGVKPKAP GVGGAFAGIP GVGPFGGPQP GVPLGYPIKA
		PKLPGGYGLP YTTGKLPYGY GPGGWAGAAG KAGYPTGTGV GPQAAAAAA KAAAKFGAGA AGVLPGVGGA

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Seq.IDNO	Name		Proteinsequenz	
			GVPGVPGAIP GIGGIAGVGT PAAAAAAAA AKAAKYGAAA GLVPGGPGFG	G PGVVGVPGAG VPGVGVPGAG
	·	٠.	IPVVPGAGIP GAAVPGVVSP EAAAKAAAKA AKYGARPGVG VGGIPTYGVG	G AGGFPGFGVG VGGIPGVAGV
			PSVGGVPGVG GVPGVGISPE AQAAAAKAA KYGVGTPAAA AAKAAAKAAQ	Q FALLNLAGLV PGVGVAPGVG
	· ·		VAPGVGVAPG VGLAPGVGVA PGVGVAPGVG VAPGIGPGGV AAAAKSAAKV	V AAKAQLRAAA GLGAGIPGLG
	` .		VGVGVPGLGV GAGVPGLGVG AGVPGFGAVP GALAAAKAAK YGAAVPGVLG	G GLGALGGVGI PGGVVGAGPA
			AAAAAAKAAA KAAQFGLVGA AGLGGLGVGG LGVPGVGGLG GIPPAAAAKA	A AKYGAAGLGG VLGGAGQFPL
			GGVAARPGFG LSPIFPGGAC LGKACGRKRK	
	01 obs - 2	Type	IV MGRDORAVAG PALRRWLLLG TVTVGFLAQS VLAGVKKFDV PCGGRDCSGG	G COCYPEKGGR GOPGPVGPOG
ית מר		٠.	YNGPPGLOGF PGLOGRKGDK	P GHPGQGGPRG RPGYDGCNGT
			_	OG LVGFQGPPGR PGHVGQMGPV
			GAPGRPGPPG PPGPKGQQGN RGLGFYGVKG EKGDVGQPGP NGIPSDTLHP	HP IIAPTGVTFH PDQYKGEKGS
	· .		EGEPGIRGIS LKGEEGIMGF PGLRGYPGLS GEKGSPGQKG SRGLDGYQGP	SP DGPRGPKGEA GDPGPPGLPA
			YSPHPSLAKG ARGDPGFPGA QCEPGSQGEP GDPGLPGPPG LSIGDGDQRR	R GLPGEMGPKG FIGDPGIPAL
	-		RGPPGPPGLP	SA RGPKGWKGDA GECRCTEGDE
			AIKGLPGLPG PKGFAGINGE PGRKGDKGDP GQHGLPGFPG LKGVPGNIGA	3A PGPKGAKGDS RTITTKGERG
			QPGVPGVPGM KGDDGSPGRD GLDGFPGLPG PPGDGIKGPP GDPGYPGIPG	PG TKGTPGEMGP PGLGLPGLKG
	· .		QRGFPGDAGL PGPPGFLGPP GPAGTPGQID CDTDVKRAVG GDRQEAIQPG	PG CIAGPKGLPG LPGPPGPTGA
-			KGLRGIPGFA GADGGPGPRG LPGDAGREGF PGPPGFIGPR GSKGAVGLPG	PG PDGSPGPIGL PGPDGPPGER
			GLPGEVLGAQ PGPRGDAGVP GQPGLKGLPG DRGPPGFRGS QGMPGMPGLK	LK GOPGLPGPSG OPGLYGPPGL
			HGFPGAPGQE GPLGLPGIPG REQLPGDRGD PGDTGAPGPV GMKGLSGDRG	RG DAGFTGEQGH PGSPGFKGID
			GMPGTPGLKG DRGSPGMDGF QGMPGLKGRP GFPGSKGEAG FFGIPGLKGL	GL AGEPGFKGSR GDPGPPGPPP

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VILDEGMEDIK PGVACPPGIT PGEKGTEGDI GLRGIRGLHG SPGLGGFPGI SGPQGRPGVF PGTVGPGRR GRPGSPGLPG FSTMPFLYCN HCPAGMRSLW BQSFQGSPSA 10 p27 MEASALTSSA MSAAAIANGG 11 Reticulocalbin MARGGRGRRL TFDQLTPDES KQATYGYYLG TLEDIDKNGD	VILPGMKDIK GEKGDEGPMG LKGYLGAKGI QGMPGIPGLS GIPGLPGRPG HIKGVKGDIG VPGIPGLPGF
p27 Reticulocalbin	SEASULEUME EASTLOANSI QOMPGIPSES SIPSEFURFO DIASVASDIG
p27 Reticulocalbin	PGVAGPPGIT GFPGFIGSRG DKGAPGRAGL YGEIGATGDF GDIGDTINLP GRPGLKGERG TTGIPGLKGF
p27 Reticulocalbin	GPPGITGVTG VQGPPGLKGQ
p27 Reticulocalbin	GLRGIRGLHG LPGTKGFPGS PGSDIHGDPG FPGPPGERGD PGEANTLPGP VGVPGQKGDQ GAPGERGPPG
p27 Reticulocalbin	SPGLQGFPGI TPPSNISGAP GDKGAPGIFG LKGYRGPPGP PGSAALPGSK GDTGNPGAPG TPGTKGWAGD
p27 Reticulocalbin	SGPQGRPGVF GLPGEKGPRG EQGFMGNTGP TGAVGDRGPK GPKGDPGFPG APGTVGAPGI AGIPQKIAIQ
p27 Reticulocalbin	PGTVGPQGRR GPPGAPGEIG PQGPPGEPGF RGAPGKAGPQ GRGGVSAVPG FRGDEGPIGH QGPIGQEGAP
p27 Reticulocalbin	GRPGSPGLPG MPGRSVSIGY LLVKHSQTDQ EPMCPVGMNK LMSGYSLLYF EGQEKAHNQD LGLAGSCLAR
p27 Reticulocalbin	FSTMPFLYCN PGDVCYYASR NDKSYWLSTT APLPMMPVAE DEIKPYISRC SVCEAPAIAI AVHSQDVSIP
p27 Reticulocalbin	HCPAGWRSLW IGYSFLMHTA AGDEGGGQSL VSPGSCLEDF RATPFIECNG GRGTCHYYAN KYSFWLTTIP
p27 Reticulocalbin	EQSFQGSPSA DTLKAGLIRT HISRCQVCMK NL
Reticulocalbin	27 MEASALTSSA VTSVAKVVRV ASGSAVVLPL ARIATVVIGG VVAMAAVPMV LSAMGFTAAG IASSSIAAKM
Reticulocalbin	MSAAAIANGG GVASGSLVGT LQSLGATGLS GLTKFILGSI GSAIAAVIAR FY
TFDQLTPDES KQATYGYYLG TLEDIDKNGD	eticulocalbin MARGGRGRRL GLALGLLLAL VLAPRVLRAK PTVRKERVVR PDSELGERPP EDNQSFQYDH EAFLGKEDSK
KQATYGYYLG TLEDIDKNGD	TFDQLTPDES KERLGKIVDR IDNDGDGFVT TEELKTWIKR VQKRYIFDNV AKVWKDYDRD KDDKISWEEY
TLEDIDKNGD	KOATYGYYLG NPAEFHDSSD HHTFKKMLPR DERRFKAADL NGDLTATREE FTAFLHPEEF EHMKEIVVLE
_	TLEDIDKNGD GFVDQDEYIA DMFSHEENGP EPDWVLSERE QFWEFRDLNK DGKLDKDEIR HWILPQDYDH
AQAEARHLVY	AQAEARHLVY ESDKNKDEKL TKEEILENWN MFVGSQATNY GEDLTKNIIDE L
12 Aldehyde MATANGAVEN	ldehyde marangaven gopdgkppal prělřnievk frkifinnem Hesksgkkfa Tcnpstreoi ceveegdkpd
dehydrogenase 6 VDKAVEAAQV	

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Abbildung 2		
Seq.IDNO	Name	Proteinsequenz
E	Gravin	LIYPRAGHADK IQGKTIPTDD NVVCFTRHEP IGVCGAITPW NFPLIAMLWHK IAPALCCGNT MULKPAGUTP LIYALYLGSLI KEAGFPPGVV NIVPGFGPTV GAAISSHEQI NKIAFTGSTE VGKLVKEAAS RSNLKRYTLE LIYALYLGSLI KEAGFPPGVV NIVPGFGPTV GAAISSHEQI NKIAFTGSTE VGKLVKEAAS RSNLKRYTLE LIGGRONFCIVC ADADLDLANE CAHGCVFRNG GOCCTAASRV FVEEQVYSEF VRRSVEYAKK R PVGDPPDVK TEGGPOIDQK QFDKILELIE SGKKEGAKLE CGGSAMEDKG LFIKPTVFSE VTDNMRIAKE EIFGPVQFIL KRKSIEEVIK RANSTDYGLT AAVFTKNLDK ALKLASALES GTVMINCYNA LYAQAPFGGF KMSGNGRELG EVALAEXTEV KTYTKLGDK NP MGAGSSTEQR SPEGPPEGSS TPAEPEPSGG GPSAEAAPDT TAAPATAASD PATKLLQKNG GLSTINGVAE ODELSLQEGD LAGQKGALAG GCALASGEEE EVIVTEVGGR BSEDVSERDS BEKNATKSAV VHDITDDGQE BENNIEDLYG AGEAASKEBE PKQSTEKPEE EVIVTEVGGR BSEDVSERDS BEKNATKSAV VHDITDDGQE BENNIEDLYG AGEAASKEBE PKQSTEKPEE TLKREGSHAE ISPPAESGGA VEECKBEGGE KOEKEBGBE KOEKEBGBE AGDHQDPSLG AGEAASKEBE PKQSTEKPEE TLKREGSHAE ISPPAESGGA VEECKBEGGE KOEKEBGBE BEPTSPYTSE TGSTFKKFFT QGÜAGURKKY SFRKPKEEDEV EASEKKKEGE PEKVDTEEDG KABVASEKLT STVEERTBEQ KTEVEETAGS VPAEELVGMD AEFQEAPPA ELVKLKETCV SGEDPTGGAD LSPDEKLEK GGESSASSPE EPEEITGLEK GLAEVQQDGE AEEGATSGE KVERGVYPMA SFKKNYTPRKK GGESSASSPE EPEEITGLEK GLAEVQQDGE AEEGATSGE KVTSVSWEA LICKGSSKR ARRBSSSDER EDELDKVKSA TLSSTESTAS BMQEBMKGSV BEPKPEEPKR KVTTSVSWEA LICKGSSKR ARRBSSSDER SKSKLLEBKSE DSIAGSGVEH STPDTEPGKE ESWVSIKKFI PGRRKKRPDG GQDPVDADENY SANISASVTEP LEQVEREABAL LTEEVLEREV RABEEPPTVT BPLPENREAR GDTVVSBARL TPEAVTARKY AANDAGTR AAABETTE WASANGLID SPDTTEBATP VQEVEGGUPD IERDENRYNG VLQAVAREKVK

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Sea IDNO	Name	Proteinsequenz
och:bac		
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		SGTQSGGTEA EAVPAQKERP PAPSSFVFQE ETKEQSKMED TLEHTDKEVS
		DEKTKDVPFF EGLEGSIDTG ITVSREKVTE VALKGEGTEE AECKKDDALE
		VOSSFASFTI, TARAEEEKVL
		TANKIIGEON
<u>.</u>		NSIPVIVE THE THE THEFT OFFICE
,		
		ALLAERIEKS LVEPKEDEKG DDVDDPENQN SALADTDASG GLTKESPDTN GPKQKEKEDA QEVELQEGNV
		HSPSDKATTP OAOEELOKOE RESAKSELTE S
		
		MANAGEBIED AWTRALLIPL LLAGPYGCLS ROELFPFGPG OGDLELEDGD DFVSPALELS GALRFYDRSD
14	Nidogen	TTAMOERDAN WEGHDGI, PPPP
-		Section of the sectio
		RNNOVPAVVA FSQGSVGFLW KSNGAYNIFA NDRESIENLA KSSNSGQQGV WVFEIGSPAT INGVVFADVI
		TOTAL TENTE TOTAL
		HQQHPQVIDV DEVELSOVI SINFNTDLHS
		PORVNGRVNG KILYGSSZYI INI

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Seq.IDNO	Name	Proteinsequenz
		VREYTMLRNI GLKTIDDIFK IAVHPLREAI DMRENMONAI VSIKELCGLP PIASLKOCLL TLSSRLITSD
		NTPSVSLVMK DSFPYLEPLG AIPDVQKKML TAYDLMIQES RFLIEMADTV QEKIVQCQKA GMEFHEELHN
		LGAKEGLKGR KLNKATESFA WNITVLKGQG DLLKNAKNEA IEWMKQIQLA CLSCGLSKAP SSSAEAKSKR
		SLEAIEEKES SEENGKL

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